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Meade, Harry
Krane, Ian

<130> 10275/041001

<141> 1999-06-15

<160> 4

<170> FastSEQ for Windows Version 4.0

<210> 1

<211> 40

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetically generated linker sequence; subsets 2 through 8 (each consisting of a repetition of the first five amino acids) encompassing positions 6 through 40 may be absent or present

<400> 1

Ser Gly Gly Gly Gly Ser Gly Gly Gly Gly Ser Gly Gly Gly Gly Ser
 1 5 10 15
 Gly Gly Gly Gly Ser Gly Gly Gly Gly Ser Gly Gly Gly Gly Ser Gly
 20 25 30
 Gly Gly Gly Ser Gly Gly Gly Gly
 35 40

<210> 2

<211> 11

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetically generated linker sequence

<400> 2

Gly Gly Gly Gly Ser Gly Gly Gly Gly Gly Ser
1 5 10

<210> 3

<211> 20

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetically generated linker sequence

<400> 1

Ser Gly Gly Gly Gly Ser Gly Gly Gly Gly Ser Gly Gly Gly Gly Ser
1 5 10 15
Gly Gly Gly Gly
20

<210> 4
<211> 17
<212> PRT
<213> Artificial Sequence

<220>
<223> Synthetically generated linker sequence

<400> 4
Ser Ser Ser Ser Gly Ser Ser Ser Ser Gly Ser Ser Ser Ser Gly Ser
1 5 10 15
Pro

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